

## Sequence Listing

## (1) GENERAL INFORMATION:

## (i) APPLICANT:

(A) NAME: Max-Planck-Gesellschaft zur Förderung der  
Wissenschaften e.V  
(B) STREET: none  
(C) CITY: Berlin  
(D) COUNTRY: Germany  
(E) POSTAL CODE: none

(ii) TITLE OF INVENTION: Process for increasing the yield in plants

(iii) NUMBER OF SEQUENCES: 1

## (iv) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk  
(B) COMPUTER: IBM PC compatible  
(C) OPERATING SYSTEM: PC-DOS/MS-DOS  
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPA)

## (2) INFORMATION FOR SEQ ID NO: 1:

## (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 1138 base pairs  
(B) TYPE: nucleotide  
(C) STRANDEDNESS: double  
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Genomic DNA

(iii) HYPOTHETICAL: NO

(iv) ANTISENSE: NO

## (vi) ORIGINAL SOURCE:

(A) ORGANISM: Agrobacterium rhizogenes

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

AATTCGATAC GAAAAAGGCA AGTGCCAGGG CCATTTAAAAA TACGGCGTCG GAAACTGGCG	60
CCAATCAGAC ACAGTCTCTG GTCCCGAAGG CCAGGGTAG TTTGGCAACA ATCACATCAA	120
GATCGATGCG CAAGACACGG GAGGCCTTAA AATCTGATC AAGCGAAAAT ACTGCATGGG	180
TGATCGTTCA CGGGTTCTATA GTACTGGGTT TGCTTTTCT TGTCGTGTTG TTGGCCCTTA	240
GGCAAAGGAT CTCAAAAGGAT GATGCCATA ATTGGGAGGA GTGGGGAAA CCTTAAAGTT	300
GGCCCGCTAT TGGATTTGGC GAAAGGGCA TTGGCAAACG TGAAGATTGC TGCATTCAAG	360
ATACCTTTTC TATTTCTGG TTAAGATGTA AAGTATTGCC ACAATCATAT TAATTACTAA	420
CATTGTATAT GAAATATACT GCGGAAATTA TCTATGCCAA AATGAATGAT TAATAATAGC	480

AATAATAATA TGTGTTAATC TTTTCAATC GGGAAATACGT CTAAGCGATT ATCGTGTGAA	540
ATAAAATTATT CCAAAAGGAA ATACATGTTT TTGGAGAACCC TGCTATAGAT ATATGCCAAA	560
TTTACACTAG TTTAGTGGGT GCAAAACATAT TATCTCTGTT TCTGAGTTA ATAAAAAAATA	580
AATAAGCAGG GCGAAATAGCA GTTACCTAA GAAGGAATGG TGCCATGTA CGTGCTTTA	600
AGAGACCTA TAATAAAATTG CCAGCTGTGT TGCTTTGGTG CCGACAGGCC TAACGTGGGG	620
TTTAGCTTGA CAAAGTAGCG CCTTCCGCA GCATAAAATAA AGGTAGGCCG GTGCCTCCCA	640
TTATTAAAGG AAAAAGCAAA AGCTGAGATT CCATAGACCA CAAACCCACCA TTATTGGAGG	660
ACAGAACCTA TTCCCTCACG TGGGTCGCTA GCTTTAAACC TAATAAGTAA AAACAATTAA	680
AAGCAGGCCAG GTGTCCCTTC TATATTCCCA CAACGAGGCCG ACGTGGACCA TCGACAGCCG	700
CATCCATTAA TTAATAAAATT TGTGGACCTA TACCTAACTC AAATATTTTT ATTATTTGCT	720
CCAATACGCT AAGAGCTCTG GATTATAAT AGTTGGATG CTTCGAGTTA TGGGGTAC	740